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TECH CENTER 1600/2900

## SEQUENCE LISTING

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<120> CYCLIC PEPTIDE LIGANDS THAT TARGET UROKINASE PLASMINOGEN  
ACTIVATOR RECEPTOR

<130> 38369-169218

<140> US 09/704,731  
<141> 2000-11-03

<150> US 09/285,783  
<151> 1999-04-05

<150> US 09/181,816  
<151> 1998-10-29

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<170> PatentIn version 3.1

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<213> Homo sapiens

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Gly Thr Cys Val Ser Asn Lys Tyr Phe Ser Asn Ile His Trp Cys Asn  
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Cys Pro Lys Lys Phe Gly Gly Gln His Cys Glu Ile Asp Lys Ser Lys  
35 40 45

Thr Cys Tyr Glu Gly Asn Gly His Phe Tyr Arg Gly Lys Ala Ser Thr  
50 55 60

Asp Thr Met Gly Arg Pro Cys Leu Pro Trp Asn Ser Ala Thr Val Leu  
65 70 75 80

Gln Gln Thr Tyr His Ala His Arg Ser Asp Ala Leu Gln Leu Gly Leu  
85 90 95

Gly Lys His Asn Tyr Cys Arg Asn Pro Asp Asn Arg Arg Arg Pro Trp  
100 105 110

Cys Tyr Val Gln Val Gly Leu Lys Leu Leu Val Gln Glu Cys Met Val  
115 120 125

His Asp Cys Ala Asp Gly Lys Lys Pro Ser Ser Pro Pro Glu Glu Leu  
130 135 140

Lys Phe Gln Cys Gly Gln Lys Thr Leu Arg Pro Arg Phe Lys Ile Ile  
 145 150 155 160  
 Gly Gly Glu Phe Thr Thr Ile Glu Asn Gln Pro Trp Phe Ala Ala Ile  
 165 170 175  
 Tyr Arg Arg His Arg Gly Gly Ser Val Thr Tyr Val Cys Gly Gly Ser  
 180 185 190  
 Leu Ile Ser Pro Cys Trp Val Ile Ser Ala Thr His Cys Phe Ile Asp  
 195 200 205  
 Tyr Pro Lys Lys Glu Asp Tyr Ile Val Tyr Leu Gly Arg Ser Arg Leu  
 210 215 220  
 Asn Ser Asn Thr Gln Gly Glu Met Lys Phe Glu Val Glu Asn Leu Ile  
 225 230 235 240  
 Leu His Lys Asp Tyr Ser Ala Asp Thr Leu Ala His His Asn Asp Ile  
 245 250 255  
 Ala Leu Leu Lys Ile Arg Ser Lys Glu Gly Arg Cys Ala Gln Pro Ser  
 260 265 270  
 Arg Thr Ile Gln Thr Ile Cys Leu Pro Ser Met Tyr Asn Asp Pro Gln  
 275 280 285  
 Phe Gly Thr Ser Cys Glu Ile Thr Gly Phe Gly Lys Glu Asn Ser Thr  
 290 295 300  
 Asp Tyr Leu Tyr Pro Glu Gln Leu Lys Met Thr Val Val Lys Leu Ile  
 305 310 315 320  
 Ser His Arg Glu Cys Gln Gln Pro His Tyr Tyr Gly Ser Glu Val Thr  
 325 330 335  
 Thr Lys Met Leu Cys Ala Ala Asp Pro Gln Trp Lys Thr Asp Ser Cys  
 340 345 350  
 Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Ser Leu Gln Gly Arg Met  
 355 360 365  
 Thr Leu Thr Gly Ile Val Ser Trp Gly Arg Gly Cys Ala Leu Lys Asp  
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 Lys Pro Gly Val Tyr Thr Arg Val Ser His Phe Leu Pro Trp Ile Arg  
 385 390 395 400  
 Ser His Thr Lys Glu Glu Asn Gly Leu Ala Leu  
 405 410

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Ile Thr Tyr Arg Phe Phe Ser Gln Ile Lys Arg  
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